OIPE

```
Input Set : N:\Crf3\RULE60\09766778.txt
                     Output Set: N:\CRF3\04262001\1766778.raw
                     SEQUENCE LISTING
      4 (1) GENERAL INFOFMATION:
             (i) APPLICANT: Papathanassiu, Adonia E
                             Green, Shawn J.
      ٠,
            (ii) TITLE OF INVENTION: Compositions and Methods for Inhibiting
     1 \odot
                                      Cellular Proliferation
     12
           (iii) NUMBER OF SEQUENCES: 2
            (iv) CORRESPONDENCE ADDRESS:
     1 4
                                                                       ENTERED
     15
                   (A) ADDRESSEE Jones & Askew
                   (B) STFEET: 191 Peachtree Street, 37th Floor
     16
     1 /
                   (C) CITY: Atlanta
     1 H
                   (D) STATE Georgia
     1 4
                   (E) COUNTRY: U.S.A.
     2^{11}
                   (F) ZIP: 30302
             (V) COMPUTER READABLE FORM
     22
                   (A) MEDIUM TYPE: Floppy disk
     23
                   (B) COMPUTER: IBM PC compatible
     24
     2 c,
                   (C) OPERATING SYSTEM: PC-DOS/MS-DOS
     24,
                   (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
     28
            (V1) CURRENT APPLICATION DATA:
C--> 29
                   (A) APPLICATION NUMBER: US/09/766,778
C--> 30
                   (B) FILING DATE: 22-Jan-2001
     31.
                   (C) CLASSIFICATION:
     3 :
           (vii) PRIOR APPLICATION DATA:
                   (A) APPLICATION NUMBER 09/227,955
     34
     35
                   (B) FILING DATE
     377
          (V111) ATTORNEY/AGENT INFOEMATION:
     313
                   (A) NAME: Greene, Jamie L.
     36
                  (B) FEGISTRATION NUMBER: 32,467
     4 (1
                   (C) PEFERENCE/DOCKET NUMBER: 05213-0290
     42
            (ix) TELECOMMUNICATION INFORMATION:
     4
                  (A) TELEPHONE: (404) 818-3700
     44
                  (B) TELEFAX: (404) 818-3799
     47
        (2) INFORMATION FOR SEQ ID NO: 1:
             (i) SEQUENCE CHARACTERISTICS:
     44
     50
                  (A) LENGTH: 276 aminc acids
    ۲. ا
                  (B) TYPE: amino acid
                  (C) STRANDEDNESS: sinule
                  (I) Torology: line at
             ii) MOLECULE TYPE: protein
     5.7
           CHIE HYPOTHETICAL NO
     59
            with ANTI-SENSE: NO
     61
             (v) FRAGMENT TYPE: N-terminal
     63
            (vi) ORIGINAL SOURCE:
     64
                  (A) ORGANISM: Homo sapiens
     f. fr
            (ix) FEATURE:
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DATE: 04/26/2001 TIME: 10:42:08

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/766,778

(A) NAME/REY: And I've site

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DATE: 04/26/2001 TIME: 10:42:08

PATENT APPLICATION: US/09/766,778

Input Set : N:\Crf3\RULE60\09766778.txt
Output Set: N:\CRF3\04262001\I766778.raw

68 (B) LOCATION: 2...3 (D) OTHER INFORMATION: /note- "Site of partial 64 70 phosphorylation." (ix) FEATURE: 7 ((A) NAME 'KEY: Active site (B) LOCATION: 117..118 7.4 (D) OTHER INFORMATION: /note= "Potential site for N-linked ٠, ٤, 76 glycosylation" ٦٠ (ix) FEATURE 74 (A) NAME (KEY: Active-site (B) LOCATION: 167..168 81 (D) OTHER INFORMATION: /note= "Potential site for N-linked 8.1 82 glycosylation" (ix) FEATURE 85 (A) NAME, KEY: Activersite (B) LOCATION: 228. 229 84. 87 (D) OTHER INFORMATION: /note= "Potential site for N-linked 8% glycosylation" 90 (ix) FEATUFE 91 (A) NAME, KEY: I-omain (B) LOCATION: 26...76 92 (D) OTHER INFORMATION: /label= Kunitz-1 94 95 (ix) FEATURE: Gt. (A) NAME/KEY: Domain (B) LOCATION: 97...147 47 (D) OTHER INFORMATION: /label= Kunitz-2 98 (ix) FEATURE: 100 (A) NAME/KEY: Domain 101 1(2)(B) LOCATION: 189..239 103 (D) OTHER INFORMATION: /label= Kunitz-3 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1 106 Asp Ser Glu Glu Asp Glu Glu His Thr Ile Ile Thr Asp Thr Glu Leu 108 109 5 10 1.5 Pro Pro Leu Lys Leu Met His Ser Phe Cys Ala Phe Lys Ala Asp Asp 1.1 10 25 30 112Gly Pro Cys Lys Ala Ile Met Lys Arg Phe Phe Phe Asn Ile Phe Thr 114 115 4011.7Arg Gln Cys Glu Glu Phe Ile Tyr Gly Gly Cys Glu Gly Asn Glr Asn 118 50 55 60 Arg Phe Glu Ser Leu Glu Glu Cys Lys Lys Met Cys Thr Arg Asp Asn 120 121 Ala Ash Aig the lie bys the The Lon Gle Gle Gle Lys Pro Asp Phe : 2 124 H^{Γ_0} dys Pho Lou dia Jim Asp Pro Gly Ile Cys Ard Gly Tyr Ile Ibr Ard 1.36 1 5 124 Tyr Pho Tur Ash Ash Gin Thr Lys Gln Cys Glu Ard Pho Lys Tyr Gly 115 130 120 132 Gly Cys Leu Gly Asn Met Asn Asn Phe Glu Thr Leu Glu Glu Cys Lys 13 + 130 135

 RAW SEQUENCE LISTING
 DATE: 04/26/2001

 PATENT APPLICATION: US/09/766,778
 TIME: 10:42:08

Input Set : N:\Crf3\RULE60\09766778.txt
Output Set: N:\CRF3\04262001\I766778.raw

135 136	Asn 145	He	Cys	Glu	Asp	Gly 150	Pro	Asn	Gly	Phe	Gln 155	Val	Asp	Asn	Tyr	Gly 160
138		Gln	Leu	Asn	Ala 165	Val	Asn	Asn	Ser	Leu 170	Thr	Pro	Gln	Ser	Thr 175	Lys
141 142	Val.	Pro	Ser	Leu 180		Glu	Phe	His	Gly 185	Pro	Ser	Trp	Cys	Leu 190	Thr	Pro
144 145	Alá	Asp	Arg 195	Gly	Leu	Cys	Arg	Ala 200	Asn	Glu	Asn	Arg	Phe 205	туг	туг	Asn
$\frac{147}{143}$	Ser	Val 210	He	31y	L/s	Суя	Arg 215	Pro	Phe	Lys	Түг	Ser 220	Gly	Cys	Gly	Gly
15.) 151	Asrı 225	Glu	Asn	Asn	Phe	Thr 230	Ser	Lys	Gln	Glu	Cys 235	Leu	Arg	Ala	Cys	Lys 240
153		Glv	Phe	He	Gln		Ile	Ser	Lys	Gly		Leu	Ile	Lys	Thr	
154	•	•			245				-	250	•			_	255	
150	Arg	Lys	Arg		L∵s	G1 r.	Arg	Val	_	Ile	Ala	Tyr	Glu		Ιlе	Phe
157		_	_	260					265					270		
15)	Val	Lys	Asn	Met												
$\frac{160}{162}$																
1 n 4	• •					refiis										
165	(-)							acids	5							
160			TYI)			o aci										
167		(C	STI	RANDI	EDNES	3S: s	sing:	lе								
168		(D) Toi	POLO	GY .	linea	ar									
170	/ i i \	(D) TOPOLOGY linear MOLECULE TYPE protein														
T 1,1	(11)	MOLI	3COPF	STY	5 E: I	prote	⊇in									
172	(iii)						∋irı									
		HYP	OTHE:	TICAI	L: NO		∋irı									
172	(iii) (iv)	HYP(OTHEN I-SEN	TICAI NSE:	L: NO			a l								
172 174	(iii) (iv)	HYPO ANT FRAC ORIO	OTHEI I-SEN GMENI GINAI	TICAI NSE: T TYI L SOU	L: NO NO PE: 1 JECE:) N-tei :	rmina									
172 174 176 178 179	(iii) (iv) (v) (vi)	HYPO ANT FRAC ORIC (A)	OTHET I-SEN GMENT GINAI () OEC	ricai NSE: r Tyl L SOU GANIS	L: NO NO PE: M JECE: SM. H) N-tei : Homo	omina sapi	iens								
172 174 176 178 179 183	(iii) (iv) (v) (vi)	HYPO ANT FRAC ORIC (A) SEQU	OTHET I-SEN GMENT GINAI) OEC JENCE	ricai NSE: r Tyl L SOU GANIS E DES	L: NO NO PE: N JECE: SM. H SCRIE	o N-tei : Home PTION	omina sap: N: SH	iens EQ II								
172 174 176 178 179 183 185	(iii) (iv) (v) (vi) (xi) Asp	HYPO ANT FRAC ORIC (A) SEQU	OTHET I-SEN GMENT GINAI) OEC JENCE	ricai NSE: r Tyl L SOU GANIS E DES	L: NO NO PE: 1 JECE: SM. H SCRIE Glu	o N-tei : Home PTION	omina sap: N: SH	iens		Asn	Ala	Glu	Ile	Cys		L⊕u
172 174 176 178 179 183 185	(iii) (iv) (v) (vi) (xi) Asp	HYPO ANT FRAC ORIC (A) SEQU Ala	OTHET I-SEN GMENT GINAI OEC JENCE Ala	FICAL NSE: F TYPE L SOU GANIS E DES	L: NO NO PE: 1 JECE: SM. H SCEIE Glu	N-ter : Home PTION Pro	omina sap: N: Si Thr	iens EQ II Gly	Asn	Asn 10				_	15	
172 174 176 178 179 183 185 186 188	(iii) (iv) (v) (vi) (xi) Asp 1	HYPO ANT FRAC ORIO (A) SEQU Ala	OTHEN	FICAL NSE: FITYEL SOU GANISE DES Gln Tyr 20	L: NO NO PE: N JECE: SM. H SCRIE Glu S	Noten : Homo PTION Pro	sapi sapi N: SI Thr Cys	iens EQ II Gly Arg	Asn Ala 25	Asn 10 Leu	Leu	Leu	Arg	Tyr 30	15 Tyr	Tyr
172 174 176 178 179 183 185 186 188	(iii) (iv) (v) (vi) (xi) Asp 1	HYPO ANT FRAC ORIO (A) SEQU Ala	OTHER I-SEN GMENT GINAL OFC JENCE Ala Asp	FICAL NSE: FITYEL SOU GANISE DES Gln Tyr 20	L: NO NO PE: N JECE: SM. H SCRIE Glu S	Noten : Homo PTION Pro	sapi sapi N: SI Thr Cys	iens EQ II Gly Arg Arg	Asn Ala 25	Asn 10 Leu	Leu	Leu	Arg Gly	Tyr 30	15 Tyr	Tyr
172 174 176 178 179 183 185 186 188 189 191	(iii) (iv) (v) (vi) (xi) Asp 1 Pro	HYPO ANT FRACORIO (A) SEQU Ala Leu	OTHEN I-SEMENT GMENT GINAL) OFC JENCE Ala Asp Tyr 35	TICAL NSE: TYPE TYPE TYPE TYPE TYPE TYPE TYPE TYPE	L: NO NO PE: I JECE: SM. H SCRIE Glu S Gly	Noter : Homo PTION Pro Pro	sapi sapi N: SH Thr Cys	iens EQ II Gly Arg Arg 40	Asn Ala 25 Gln	Asn 10 Leu Phe	Leu Leu	Leu Tyr	Arg Gly 45	Tyr 30 Gly	15 Tyr Cys	Tyr Glu
172 174 176 178 179 183 185 186 188 191 191	(iii) (iv) (v) (vi) (xi) Asp 1 Pro	HYPO ANT FRAC OEIO (A) SEQU Ala Leu Ang	OTHEN I-SEMENT GMENT GINAL) OFC JENCE Ala Asp Tyr 35	TICAL NSE: TYPE TYPE TYPE TYPE TYPE TYPE TYPE TYPE	L: NO NO PE: I JECE: SM. H SCRIE Glu S Gly	Noter : Homo PTION Pro Pro	sapi sapi N: SH Thr Cys	iens EQ II Gly Arg Arg	Asn Ala 25 Gln	Asn 10 Leu Phe	Leu Leu	Leu Tyr	Arg Gly 45	Tyr 30 Gly	15 Tyr Cys	Tyr Glu
172 174 176 178 179 183 185 186 188 189 191	(iii) (iv) (v) (vi) (xi) Asp 1 Pro Asp	HYPO ANT FRAC ORIC (A) SEQU Ala Leu Arg Asn 50	OTHER I-SEN I-SEN ISINAI OFC JENCE Ala Asp Tyr 35 Ala	TICAN NSE: TIYH L SOU GANIS E DES Gln Tyr 20 Thr	NO N	N-ten : Homo PTION Pro Pro Ser	sapi Sapi N: SH Thr Cys Cys Tyr	iens EQ II Gly Arg Arg 40 Thr	Asn Ala 25 Gln Trp	Asn 10 Leu Phe Glu	Leu Leu Ala	Leu Tyr Cys 60	Arg Gly 45 Asp	Tyr 30 Gly Asp	15 Tyr Cys Ala	Tyr Glu Cys
172 174 176 178 179 183 185 186 188 191 192 194 195	(iii) (iv) (v) (vi) (xi) Asp 1 Pro Asp	HYPO ANT FRAC ORIC (A) SEQU Ala Leu Arg Asn 50	OTHER I-SEN I-SEN ISINAI OFC JENCE Ala Asp Tyr 35 Ala	TICAN NSE: TIYH L SOU GANIS E DES Gln Tyr 20 Thr	NO N	N-ten : Homo PTION Pro Pro Ser	sapi Sapi N: SH Thr Cys Cys Tyr	iens EQ II Gly Arg Arg 40	Asn Ala 25 Gln Trp	Asn 10 Leu Phe Glu	Leu Leu Ala	Leu Tyr Cys 60	Arg Gly 45 Asp	Tyr 30 Gly Asp	15 Tyr Cys Ala	Tyr Glu Cys
172 174 176 178 179 183 186 186 189 191 192 194 195	(iii) (iv) (v) (vi) (xi) Asp 1 Pro Asp Gly Trp 65	HYPO ANT FRAC ORIC (A) SEQU Ala Leu Arg Asn 50 Arg	OTHER I-SEN I-SEN IMMENT GINAI) OFC JENCE Ala Asp Tyr 35 Ala Ile	TICANSE: TYNSE: TYNSEANISE TYNSEANISE TYN 20 Thr Asn	L: NO NO PE: 1 JECE: SM. H GCHIH Glu S Gly Gln Asn Lys	Noter: Homo PTION Pro Pro Ser Phe Val	sapi Sapi N: Sh Thr Cys Cys Tyr 55 Pro	iens EQ II Gly Arg Arg 40 Thr	Asn Ala 25 Gln Trp Val	Asn 10 Leu Phe Glu Cys	Leu Leu Ala Arg	Leu Tyr Cys 60 Leu	Arg Gly 45 Asp Gln	Tyr 30 Gly Asp	15 Tyr Cys Ala Ser	Tyr Glu Cys Val 80
172 174 176 178 179 183 185 186 188 191 192 194 195 197 198	(iii) (iv) (v) (vi) (xi) Asp 1 Pro Asp Gly Trp 65	HYPO ANT FRAC ORIC (A) SEQU Ala Leu Arg Asn 50 Arg	OTHER I-SEN I-SEN IMMENT GINAI) OFC JENCE Ala Asp Tyr 35 Ala Ile	TICANSE: TYNSE: TYNSEANISE TYNSEANISE TYN 20 Thr Asn	L: NO NO PE: 1 JECE: SM. H GCHIH Glu S Gly Gln Asn Lys	Noter: Homo PTION Pro Pro Ser Phe Val	sapi Sapi N: Sh Thr Cys Cys Tyr 55 Pro	iens EQ II Gly Arg Arg 40 Thr	Asn Ala 25 Gln Trp Val	Asn 10 Leu Phe Glu Cys	Leu Leu Ala Arg 75 Tyr	Leu Tyr Cys 60 Leu	Arg Gly 45 Asp Gln The	Tyr 30 Gly Asp Val	15 Tyr Cys Ala Ser Leu	Tyr Glu Cys Val 80 Ser
172 174 176 178 179 183 185 186 188 191 192 194 195 197 198	(iii) (iv) (v) (vi) (xi) Asp 1 Pro Asp Gly Trp 65	HYPO ANT FRAC ORIC (A) SEQU Ala Leu Arg Asn 50 Arg	OTHER I-SEN GMENT GINAL) OFC JENCE Ala Asp Tyr 35 Ala Ile	TICANSE: TYIL SOUD SEARCH TYIL SOUD SEARCH TYIL SOUD TYIL SOUD TYIL TYIL TYIL TYIL TYIL TYIL TYIL TYIL	L: NO	Note: Homo PTION Pro Pro Pro Ser Phe Val Gly	sap: N: SH Thr Cys Cys Tyr 55 Pro	iens EQ II Gly Arg Arg 40 Thr	Asn Ala 25 Gln Trp Val	Asn 10 Leu Phe Glu Cys	Leu Leu Ala Arg 75 Tyr	Leu Tyr Cys 60 Leu	Arg Gly 45 Asp Gln	Tyr 30 Gly Asp Val	15 Tyr Cys Ala Ser Leu	Tyr Glu Cys Val 80 Ser
172 174 176 178 179 183 185 186 189 191 192 194 195 197 198 201 201 201 4	(iii) (iv) (v) (vi) (xi) Asp 1 Pro Asp Gly Trp 65 Asp	HYPCANT FRACORIO (A) SEQUALA Ala Leu Arg Asn 50 Arg Asp.	OTHER I-SEN GMENT GINAL) OFC JENCE Ala Asp Tyr 35 Ala Ile G.n	TICANSE: TYME SOUTH TY SOUTH TYME SOUTH TYME SOUTH TYME SOUTH TYME SOUTH TYME SOUTH TYNE SOUTH TYME SOUTH TYME SOUTH TY	L: NO NO PE: M PECE: M	Note: Homo PTION Pro Pro Ser Phe Val Giy	sap: Sap: SI Thr Cys Cys Tyr 55 Pro Ser	iens EQ II Gly Arg Arg 40 Thr Lys	Asn Ala 25 Gln Trp Val Gla Ser L	Asn 10 Leu Phe Glu Cys Mys	Leu Leu Ala Arg 75 Tyr	Leu Tyr Cys 60 Leu The	Arg Gly 45 Asp Gln The	Tyr 30 Gly Asp Val Asn Ard	15 Tyr Cys Ala Ser Len 95 Asn	Tyr Glu Cys Val 80 Ser Ara
172 174 176 178 179 183 185 186 189 191 192 194 195 197 198 201 201 206	(iii) (iv) (v) (vi) (xi) Asp 1 Pro Asp Gly Trp 65 Asp	HYPCANT FRACORIO (A) SEQUALA Ala Leu Arg Asn 50 Arg Asp.	OTHER I-SEN GMENT GINAL) OFC JENCE Ala Asp Tyr 35 Ala Ile G.n Thr	TICANSE: TYME SOUTH TY SOUTH TYME SOUTH TYME SOUTH TYME SOUTH TYME SOUTH TYME SOUTH TYNE SOUTH TYME SOUTH TYME SOUTH TY	L: NO NO PE: M PECE: M	Note: Homo PTION Pro Pro Ser Phe Val Giy	sap: Sap: SI Thr Cys Cys Tyr 55 Pro Ser	iens EQ II Gly Arg Arg 40 Thr Lys Ill:	Asn Ala 25 Gln Trp Val Gla Ser L	Asn 10 Leu Phe Glu Cys Mys	Leu Leu Ala Arg 75 Tyr	Leu Tyr Cys 60 Leu The	Arg Gly 45 Asp Gln The His	Tyr 30 Gly Asp Val Asn Ard	15 Tyr Cys Ala Ser Len 95 Asn	Tyr Glu Cys Val 80 Ser Ara
172 174 176 178 179 183 185 186 189 191 192 194 195 197 198 201 204 206 207	(iii) (iv) (v) (vi) (xi) Asp 1 Pro Asp Gly Trp 65 Asp Ser	HYPE ANT FRAC ORIO (A SEQU Ala Leu Arg Asn 50 Arg Asp Med	OTHER I-SEN GMENT GINAL) OFC JENCE Ala Asp Tyr 35 Ala Ile G.n Thr Asn 1:5	TICANSE: TYME SOUTH TY SOUTH TYME SOUTH TYME SOUTH TYME SOUTH TYME SOUTH TYME SOUTH TYNE SOUTH TYME SOUTH TY S	L: NO NO PE: NO PE: M JECE: SM. H SCHIE Glu S Gly Gin Asn Lys Cla E5	N-ter: Homo PTION Pro Pro Pro Ser Phe Val 70 Gly Lys	sap: Sap: SI Thr Cys Cys Tyr 55 Pro Ser Pho	iens EQ II Gly Arg Arg 40 Thr Lys Flo	Asn Ala 25 Gln Trp Val Gls Ser 1 f Ala	Asn 10 Leu Phe Glu Cys Mys My Thr	Leu Leu Ala Arg 75 Tyr Gys	Leu Tyr Cys 60 Leu The Cys Met	Arg Gly 45 Asp Gln The His	Tyr 30 Gly Asp Val Ass Arg	15 Tyr Cys Ala Ser Leu 95 Asn	Tyr Glu Cys Val 80 Ser Ard
172 174 176 178 179 183 185 186 189 191 192 194 195 197 198 201 201 206	(iii) (iv) (v) (vi) (xi) Asp 1 Pro Asp Gly Trp 65 Asp Ser	HYPE ANT FRAC ORIO (A SEQU Ala Leu Arg Asn 50 Arg Asp Med	OTHER I-SEN GMENT GINAL) OFC JENCE Ala Asp Tyr 35 Ala Ile G.n Thr Asn 1:5	TICANSE: TYME SOUTH TY SOUTH TYME SOUTH TYME SOUTH TYME SOUTH TYME SOUTH TYME SOUTH TYNE SOUTH TYME SOUTH TY S	L: NO NO PE: NO PE: M JECE: SM. H SCHIE Glu S Gly Gin Asn Lys Cla E5	N-ter: Homo PTION Pro Pro Pro Ser Phe Val 70 Gly Lys	sap: Sap: SI Thr Cys Cys Tyr 55 Pro Ser Pho	iens EQ II Gly Arg Arg 40 Thr Lys Ill:	Asn Ala 25 Gln Trp Val Gls Ser 1 f Ala	Asn 10 Leu Phe Glu Cys Mys My Thr	Leu Leu Ala Arg 75 Tyr Gys	Leu Tyr Cys 60 Leu The Cys Met	Arg Gly 45 Asp Gln The His	Tyr 30 Gly Asp Val Ass Arg	15 Tyr Cys Ala Ser Leu 95 Asn	Tyr Glu Cys Val 80 Ser Ard

RAW SEQUENCE LISTING

DATE: 04/26/2001

PATENT APPLICATION: US/09/766,778 TIME: 10:42:08

Input Set : N:\Crf3\RULE60\09766778.txt
Output Set: N:\CRF3\04262001\1766778.raw

212	Cys	Ser	Ala	Asn	Val	Thr	Arg	туг	туг	Phe	Asn	Pro	Arg	Tyr	Arg	Thr
213	145					150					155					160
215	Cys	Asp	Ala	Phe	Thr	lyr	Thr	Gly	Cys	Gly	Gly	Asn	Asp	Asn	Asn	Phe
216					165					170					175	
218	Val	Ser	Arg	Glu	Asp	Cys	Lys	Arg	Ala	Cys	Ala	Lys	Ala	Leu	Lys	Lys
219				180					185					190		
121	Lys	Lys	Lys	Met	Pro	Lys	Leu	Arg	Phe	Ala	Ser	Arg	Пlе	Arg	Lys	lle
222			195					200					205			
224	Arg	Lys	Lys	Gln	Phe											
225		210														

VERIFICATION SUMMARY

DATE: 04/26/2001 TIME: 10:42:09

PATENT APPLICATION: US/09/766,778

Input Set : N:\Crf3\RULE60\09766778.txt Output Set: N:\CRF3\04262001\1766778.raw

L:29 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]

L:30 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]